PCT

10/522341

PTO/SB/92 (09-04)
Approved for use through 7/31/2006. OMB 0651-0031
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#4

Application No. (if known): 10/522341

Attorney Docket No.: 12810-00057-US

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Amendment and Reply to Notification to Comply with Requirements for Patent Applications Containing Nucleotide and/or Amino Acid Sequence Disclosures (9 pages)
Copy of Notification to Comply With Requirements for Patent Applications Containing Nucleotide and/or Amino Acid Sequence Disclosures (12 pages)
Statement to Support Filing and Submission of Substitute Sequence Listing in Accordance with 37 CFR §§1.821 through 1.825 (2 pages)
Replacement Nucleotide and/or Amino Acid Sequence Submission on CD (COPY 1 REPLACEMENT 03/03/2006, COPY 2 REPLACEMENT 03/03/2006, CRF COPY REPLACEMENT 03/03/2006, 3 total discs)

Certificate of Mailing (1 page)
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U.S. APPLICATION NUMBER NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
10/522,341	Michael Kock	532622010400

INTERNATIONAL APPLICATION NO.	
PCT/EP03/07877	
I.A. FILING DATE	PRIORITY DATE
07/18/2003	07/26/2002

23416
CONNOLLY BOVE LODGE & HUTZ, LLP
P O BOX 2207
WILMINGTON, DE 19899

CONFIRMATION NO. 5941

371 FORMALITIES LETTER

OC000000017724246

Date Mailed: 01/03/2006

NOTIFICATION TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant is given **TWO MONTHS FROM THE DATE OF THIS NOTICE** within which to file the items indicated below to avoid abandonment. Extensions of time may be obtained under the provisions of 37 CFR 1.136(a).

- A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

Applicant is cautioned that correction of the above items may cause the specification and drawings page count to exceed 100 pages. If the specification and drawings exceed 100 pages, applicant will need to submit the required application size fee.

For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:

- **For Rules Interpretation, call (571) 272-0951**
- **For Patentin Software Program Help, call Patent EBC at 1-866-217-9197 or directly at 703-305-3028 / 703-308-6845 between the hours of 6 a.m. and 12 midnight, Monday through Friday, EST.**
- **Send e-mail correspondence for Patentin Software Program Help @ ebc@uspto.gov**

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

*A copy of this notice **MUST** be returned with the response.*

WINSTON M ALVARADO

Telephone: (703) 308-9140 EXT 206

PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
10/522,341	PCT/EP03/07877	532622010400

FORM PCT/DO/EO/922 (371 Formalities Notice)

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/522,341
Source: PCT
Date Processed by STIC: 2-17-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.2.2 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):** U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

Need mail flag report with the notification of missing parts

Allison Ahrendt
National Stage Processing
Patents Specialist
(703) 305-3421

PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/522,341

DATE: 02/17/2005
TIME: 12:04:43

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Output Set: N:\CRF4\02172005\J522341.raw

2 <110> APPLICANT: Kock, Michael
3 Frank, Markus
4 Badur, Ralf
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8 <130> FILE REFERENCE: 532622010400
C > 10 <140> CURRENT APPLICATION NUMBER: US/10/522,341
C > 11 <141> CURRENT FILING DATE: 2005-01-25
13 <160> NUMBER OF SEQ ID NOS: 71
15 <170> SOFTWARE: PatentIn Ver. 2.1

ORED SEQUENCES

(pg. 1-4, 6)

350 <210> SEQ ID NO: 4
351 <211> LENGTH: 427
352 <212> TYPE: PRT
353 <213> ORGANISM: Artificial sequence
W > 354 <220> FEATURE: — pls insert
354 <223> OTHER INFORMATION: Description of the artificial sequence: coding for
355 cytosine deaminase (codA)
E > 357 <400> SEQUENCE: 4
358 Met Ser Asn Asn Ala Leu Gln Thr Ile Ile Asn Ala Arg Leu Pro Gly
359 1 5 10 15
361 Glu Glu Gly Leu Trp Gln Ile His Leu Gln Asp Gly Lys Ile Ser Ala
362 20 25 30
364 Ile Asp Ala Gln Ser Gly Val Met Pro Ile Thr Glu Asn Ser Leu Asp
365 35 40 45
367 Ala Glu Gln Gly Leu Val Ile Pro Pro Phe Val Glu Pro His Ile His
368 50 55 60
370 Leu Asp Thr Thr Gln Thr Ala Gly Gln Pro Asn Trp Asn Gln Ser Gly
371 65 70 75 80
373 Thr Leu Phe Glu Gly Ile Glu Arg Trp Ala Glu Arg Lys Ala Leu Leu
374 85 90 95
376 Thr His Asp Asp Val Lys Gln Arg Ala Trp Gln Thr Leu Lys Trp Gln
377 100 105 110
379 Ile Ala Asn Gly Ile Gln His Val Arg Thr His Val Asp Val Ser Asp
380 115 120 125
382 Ala Thr Leu Thr Ala Leu Lys Ala Met Leu Glu Val Lys Gln Glu Val
383 130 135 140
385 Ala Pro Trp Ile Asp Leu Gln Ile Val Ala Phe Pro Gln Glu Gly Ile
386 145 150 155 160
388 Leu Ser Tyr Pro Asn Gly Glu Ala Leu Leu Glu Glu Ala Leu Arg Leu
389 165 170 175

FYI: pls insert <220> whenever <221>, <222> or <223> is present.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/522,341

DATE: 02/17/2005
TIME: 12:04:43

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Output Set: N:\CRF4\02172005\J522341.raw

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394 Tyr Gly Val Glu Ser Leu His Lys Thr Phe Ala Leu Ala Gln Lys Tyr
395         195                 200                 205
397 Asp Arg Leu Ile Asp Val His Cys Asp Glu Ile Asp Asp Glu Gln Ser
398     210                 215                 220
400 Arg Phe Val Glu Thr Val Ala Ala Leu Ala His His Glu Gly Met Gly
401 225                 230                 235                 240
403 Ala Arg Val Thr Ala Ser His Thr Thr Ala Met His Ser Tyr Asn Gly
404                 245                 250                 255
406 Ala Tyr Thr Ser Arg Leu Phe Arg Leu Leu Lys Met Ser Gly Ile Asn
407             260                 265                 270
409 Phe Val Ala Asn Pro Leu Val Asn Ile His Leu Gln Gly Arg Phe Asp
410         275                 280                 285
412 Thr Tyr Pro Lys Arg Arg Gly Ile Thr Arg Val Lys Glu Met Leu Glu
413     290                 295                 300
415 Ser Gly Ile Asn Val Cys Phe Gly His Asp Asp Val Phe Asp Pro Trp
416 305                 310                 315                 320
418 Tyr Pro Leu Gly Thr Ala Asn Met Leu Gln Val Leu His Met Gly Leu
419                 325                 330                 335
421 His Val Cys Gln Leu Met Gly Tyr Gly Gln Ile Asn Asp Gly Leu Asn
422             340                 345                 350
424 Leu Ile Thr His His Ser Ala Arg Thr Leu Asn Leu Gln Asp Tyr Gly
425         355                 360                 365
427 Ile Ala Ala Gly Asn Ser Ala Asn Leu Ile Ile Leu Pro Ala Glu Asn
428     370                 375                 380
430 Gly Phe Asp Ala Leu Arg Arg Gln Val Pro Val Arg Tyr Ser Val Arg
431 385                 390                 395                 400
433 Gly Gly Lys Val Ile Ala Ser Thr Gln Pro Ala Gln Thr Thr Val Tyr
434                 405                 410                 415
436 Leu Glu Gln Pro Glu Ala Ile Asp Tyr Lys Arg
437             420                 425
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4799 <211> LENGTH: 27
4800 <212> TYPE: DNA
4801 <213> ORGANISM: Artificial sequence
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4803 <223> OTHER INFORMATION: Description of the artificial sequence:
4804       oligonucleotide primer
4806 <400> SEQUENCE: 50
W--> 4807 cgtgaatacg gcgtggagtc g
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4813 <213> ORGANISM: Artificial sequence
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4816       oligonucleotide primer
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Found 21

Found 20

21 OK

RAW SEQUENCE LISTING DATE: 02/17/2005
PATENT APPLICATION: US/10/522,341 TIME: 12:04:44

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Output Set: N:\CRF4\02172005\J522341.raw

(20) - see pg. 2 For error explanation

E--> 4819 cggcaggata atcaggttgg
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4843 <211> LENGTH: (27) - Found 20
4844 <212> TYPE: DNA
4845 <213> ORGANISM: Artificial sequence
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4847 <223> OTHER INFORMATION: Description of the artificial sequence:
4848 oligonucleotide primer
4850 <400> SEQUENCE: 53
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5507 <211> LENGTH: 1011
5508 <212> TYPE: DNA
5509 <213> ORGANISM: Zea mays
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5511 <221> NAME/KEY: CDS
5512 <222> LOCATION: (1)..(981)
5513 <223> OTHER INFORMATION: coding for 5-methylthioribose kinase
5515 <400> SEQUENCE: 59

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5517 Ala Arg Ala Leu Leu Ser Ser Pro Leu Ala Gly Ala Ser Pro Asp Cys
5518   1               5                  10                  15
5520 cag tca gcc tca gcc atg gcc gcg gag gag gag cag ggc ttc cgc ccg   96
5521 Gln Ser Ala Ser Ala Met Ala Ala Glu Glu Glu Gln Gly Phe Arg Pro
5522             20                  25                  30
5524 ctg gac gag tcg tcc ctg ctc gcc tac atc aag gcc acg ccg gcg ctc  144
5525 Leu Asp Glu Ser Ser Leu Leu Ala Tyr Ile Lys Ala Thr Pro Ala Leu
5526          35                  40                  45
5528 gcc tcc cgc ctc ggc ggc ggt ggc agt cta gac tcc atc gag atc aag  192
5529 Ala Ser Arg Leu Gly Gly Gly Gly Ser Leu Asp Ser Ile Glu Ile Lys
5530      50                  55                  60
5532 gag gtc ggc gac ggc aac ctc aac ttc gtc tac atc gtg cag tcc gag  240
5533 Glu Val Gly Asp Gly Asn Leu Asn Phe Val Tyr Ile Val Gln Ser Glu
5534  65                  70                  75                  80
5536 gcc ggc gcc atc gtc gtc aag cag gcg ctc ccg tac gtg cgc tgc gtg  288
5537 Ala Gly Ala Ile Val Val Lys Gln Ala Leu Pro Tyr Val Arg Cys Val
5538                  85                  90                  95
5540 ggg gat tcg tgg ccc atg acg cgg gag cgc gcc tac ttc gag gcc tcc  336
5541 Gly Asp Ser Trp Pro Met Thr Arg Glu Arg Ala Tyr Phe Glu Ala Ser
5542             100                 105                 110
5544 acg ctg cgg gag cac ggc cgc ctg tgc ccg gag cac acc ccc gag gtg  384
5545 Thr Leu Arg Glu His Gly Arg Leu Cys Pro Glu His Thr Pro Glu Val
5546         115                 120                 125
5548 tac cac ttc gac cgg acc ttg tcg ctg atg ggg atg cgc tac atc gag  432
5549 Tyr His Phe Asp Arg Thr Leu Ser Leu Met Gly Met Arg Tyr Ile Glu
5550     130                 135                 140
5552 ccc ccg cac atc atc ctc cgc aag ggc ctc gtc gcc ggt gtc gag tac  480
5553 Pro Pro His Ile Ile Leu Arg Lys Gly Leu Val Ala Gly Val Glu Tyr
5554 145                 150                 155                 160
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RAW SEQUENCE LISTING DATE: 02/17/2005
PATENT APPLICATION: US/10/522,341 TIME: 12:04:44

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\02172005\J522341.raw

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5558                 165                 170                 175
5560 ttc acc tcc ctc ctc tat aac aat acc acg gat cat aag aac gga gtt   576
5561 Phe Thr Ser Leu Leu Tyr Asn Asn Thr Thr Asp His Lys Asn Gly Val
5562             180                 185                 190
5564 gct aag tac tct gcg aac gtg gag atg tgt agg ctc acg gag caa gtt   624
5565 Ala Lys Tyr Ser Ala Asn Val Glu Met Cys Arg Leu Thr Glu Gln Val
5566         195                 200                 205
5568 gtg ttc tcg gac cca tac cgt gtt tcc aaa ttt aat cgg tgg acc tcg   672
5569 Val Phe Ser Asp Pro Tyr Arg Val Ser Lys Phe Asn Arg Trp Thr Ser
5570     210                 215                 220
5572 cct tat ctc gac aaa gat gct gag gca gtt cgc gag gat gat gag ctc   720
5573 Pro Tyr Leu Asp Lys Asp Ala Glu Ala Val Arg Glu Asp Asp Glu Leu
5574 225                 230                 235                 240
5576 aag ttg gaa gta gct ggg ctg aaa tcg atg ttt atc gag aga gct caa   768
5577 Lys Leu Glu Val Ala Gly Leu Lys Ser Met Phe Ile Glu Arg Ala Gln
5578                 245                 250                 255
5580 gct ctg att cat gga gat ctc cac act ggt tct atc atg gtg acc gaa   816
5581 Ala Leu Ile His Gly Asp Leu His Thr Gly Ser Ile Met Val Thr Glu
5582             260                 265                 270
5584 gtt caa ctc aag tca ttg atc cag aat ttg ggt tct atg ggg cca atg   864
5585 Val Gln Leu Lys Ser Leu Ile Gln Asn Leu Gly Ser Met Gly Pro Met
5586         275                 280                 285
5588 ggg ttt gat att ggg agc ctt cct tgg aaa cct gat ttt ggg cat act   912
5589 Gly Phe Asp Ile Gly Ser Leu Pro Trp Lys Pro Asp Phe Gly His Thr
5590     290                 295                 300
5592 atg cac aga atg ggc atg ctg atc aag cga atg atc gta agg ctt aca   960
5593 Met His Arg Met Gly Met Leu Ile Lys Arg Met Ile Val Arg Leu Thr
5594 305                 310                 315                 320
5596 aga atg gat ctt gaa gac aat tgaagagtcg tggaatttgt tccacaaaaa      1011
E--> 5597 arg met asp Leu glu asp asn                          325
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325

move to 325 delete

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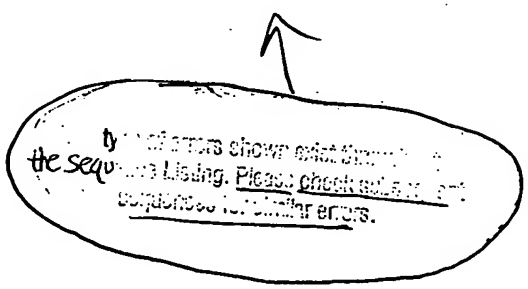
tataataatt atcattaatt agtagtaata taatatttca aatatttttt tcaaaataaa 4800
agaatgtagt atatagcaat tgcttttctg tagtttataa gtgtgtatat tttaatttat 4860
aacttttcta atatatgacc aaaatttgtt gatgtgcagg tatcaccgga tccatcgaat 4920
tcggtacgct gaaatcacca gtctctctct acaaatctat ctctctctat tttctccata 4980
aataatgtgt gagtagtttc ccgataaggg gaantttaggg ttcttatagg gtttcgctca 5040
tgtgttgagc atataagaaa cccttagtat gtatttgtat ttgtaaaata cttctatcaa 5100
taaaatttct aattcctaaa accaaaatcc agtactaaaa tccagatctc ctaaagtccc 5160
tatagatctt tgtcgtgaat ataaaccaga cacgagacga ctaaacctgg agcccagacg 5220
ccgttcgaag ctagaagtac cgcttaggca ggaggccgtt agggaaaaga tgctaaggca 5280
gggttggtta cgttgactcc cccgtaggtt tggtttaaat atgatgaagt ggacggaagg 5340
aaggaggaag acaaggaagg ataaggttgc aggccctgtg caaggtaaga agatggaaat 5400
ttgatagagg tacgctacta tacttatact atacgctaag ggaatgcttg tatttatacc 5460
ctataccccc taataacccc ttatcaattt aagaaataat ccgcataagc ccccgcttaa 5520
aaattggtat cagagccatg aataggtcta tgaccaaaac tcaagaggat aaaacctcac 5580
caaaatacga aagagttctt aactctaaag ataaaagatc tttcaagatc aaaactagtt 5640
ccctcacacc ggtgacgggg atcgcgatgg gtac 5674

Pls explain "N" location.

See error explanation on page 7.



VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/10/522,341

DATE: 02/17/2005
TIME: 12:04:45

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\02172005\J522341.raw

Error Explanation: 2

Use of n's or Xaa's (NEW RULES):
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:55; N Pos. 5014
Seq#:57; N Pos. 6697
Seq#:59; N Pos. 1026

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/522,341

DATE: 02/17/2005
TIME: 12:04:45

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\02172005\J522341.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:228 M:283 W: Missing Blank Line separator, <220> field identifier
L:231 M:283 W: Missing Blank Line separator, <220> field identifier
L:236 M:283 W: Missing Blank Line separator, <220> field identifier
L:354 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:357 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:4
L:443 M:283 W: Missing Blank Line separator, <220> field identifier
L:641 M:283 W: Missing Blank Line separator, <220> field identifier
L:645 M:283 W: Missing Blank Line separator, <400> field identifier
L:866 M:283 W: Missing Blank Line separator, <220> field identifier
L:1092 M:283 W: Missing Blank Line separator, <220> field identifier
L:1199 M:283 W: Missing Blank Line separator, <220> field identifier
L:1384 M:283 W: Missing Blank Line separator, <220> field identifier
L:1568 M:283 W: Missing Blank Line separator, <220> field identifier
L:1713 M:283 W: Missing Blank Line separator, <220> field identifier
L:1802 M:283 W: Missing Blank Line separator, <220> field identifier
L:1891 M:283 W: Missing Blank Line separator, <220> field identifier
L:2014 M:283 W: Missing Blank Line separator, <220> field identifier
L:2260 M:283 W: Missing Blank Line separator, <220> field identifier
L:2606 M:283 W: Missing Blank Line separator, <220> field identifier
L:2834 M:283 W: Missing Blank Line separator, <220> field identifier
L:3185 M:283 W: Missing Blank Line separator, <220> field identifier
L:3413 M:283 W: Missing Blank Line separator, <220> field identifier
L:3642 M:283 W: Missing Blank Line separator, <220> field identifier
L:3754 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:37
L:3847 M:283 W: Missing Blank Line separator, <220> field identifier
L:4039 M:283 W: Missing Blank Line separator, <220> field identifier
L:4225 M:283 W: Missing Blank Line separator, <220> field identifier
L:4411 M:283 W: Missing Blank Line separator, <220> field identifier
L:4597 M:283 W: Missing Blank Line separator, <220> field identifier
L:4782 M:283 W: Missing Blank Line separator, <220> field identifier
L:4802 M:283 W: Missing Blank Line separator, <220> field identifier
L:4807 M:252 E: No. of Seq. differs, <211> LENGTH:Input:27 Found:21 SEQ:50
L:4814 M:283 W: Missing Blank Line separator, <220> field identifier
L:4819 M:252 E: No. of Seq. differs, <211> LENGTH:Input:26 Found:20 SEQ:51
L:4826 M:283 W: Missing Blank Line separator, <220> field identifier
L:4846 M:283 W: Missing Blank Line separator, <220> field identifier
L:4851 M:252 E: No. of Seq. differs, <211> LENGTH:Input:27 Found:20 SEQ:53
L:4857 M:283 W: Missing Blank Line separator, <220> field identifier
L:4869 M:283 W: Missing Blank Line separator, <220> field identifier
L:4957 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:55
L:4957 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:55
L:4957 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:4980
L:4975 M:283 W: Missing Blank Line separator, <220> field identifier
L:5087 M:283 W: Missing Blank Line separator, <220> field identifier
L:5203 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:57
L:5203 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:57

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/522,341

DATE: 02/17/2005
TIME: 12:04:45

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\02172005\J522341.raw

L:5203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:6660
L:5262 M:283 W: Missing Blank Line separator, <220> field identifier
L:5510 M:283 W: Missing Blank Line separator, <220> field identifier
L:5597 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:1011
L:5597 M:254 E: No. of Bases conflict, LENGTH:Input:325 Counted:1032 SEQ:59
L:5597 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6
L:5597 M:112 C: (48) String data converted to lower case,
L:5597 M:252 E: No. of Seq. differs, <211> LENGTH:Input:1011 Found:1032 SEQ:59
L:5673 M:283 W: Missing Blank Line separator, <220> field identifier
L:5761 M:283 W: Missing Blank Line separator, <220> field identifier
L:5842 M:283 W: Missing Blank Line separator, <220> field identifier
L:5923 M:283 W: Missing Blank Line separator, <220> field identifier
L:6003 M:283 W: Missing Blank Line separator, <220> field identifier
L:6015 M:283 W: Missing Blank Line separator, <220> field identifier
L:6027 M:283 W: Missing Blank Line separator, <220> field identifier